

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P40511	SPO22_YEAST	Saccharomyces cerevisiae	Sporulation-specific protein 22	24.475439	NaN	NaN	33229814	MSDHNVNSTFRKTLVELCETATWIT SQVYAAKNLEKNDLITVDNKISALYP IAEKYDRSFRITTVILDEELILKLENA ASSLWNSLTIAMKAEKASDKYFNEV FCKCKIFATKLLSIHEALFRNTNLL RNFKCYISSFKSASEYRFDDLITNTQ QHSEKYLQIINENVESFSNEEKTEFK KLTFEFYLVNFQLYLSENDLDTANIY TAKVNITDNSKYMDADLLIELCRMIY NSTVMLKEINNPETQLVDVNIISFLK DVEKYLELPVENLKSHTDYSNLKYS VLIFMANCLVEGHPQASELEQCDHY LSLLQNEYPNKVDPFILAINLTKRRN IVNPAETIEEILMRMIMSVDVISNFO AVIASINDLSKMNTKFSIVCLDYLLIN KLNSKNDSKFLGKAICSRFLITTSK TMNDSEIAESLENFSTQMERIVSEP LTKHAISCITLLWNTGKKLEKMEKY VVSIRFYKLALKDIISQNYSDRGKIQR ALQVVYNKIEDYSNTVRVYQDMDEV DROSPLCQLLMLQSFLLADDKTEEAL TCLQKIKSSEDEKSTDALILAVAECK RKTDLQSVQGLLMIFDKLQSKSNSQTI SSTSSSQTLILRYTLQMIKVSSEEP LETFINYLPTVQKLLQKAVEFLKTVK LLNQLPPDVEKEAIYQOSVAVNEIE WFASFSYNVAVKCLVDQSCESISSEF PQYCIQFIDLIPVQDFTFPKMYHFTY WRFKATILQLIAKEKAKQDQHOKD WDIYEKSEELVNSINVMKKSSEFKD GSSLEDRNTLHECFLEALTIHLESAL MMPDQTRILDILKKTLEYQDSRVDA LLIDISSNMEDLPKGVLEIILETVLKR NMGPEVKERELCSWLRILLENAINL NHEVELRILDRVLKILNINQSSLQDT DGVLQTELETIATYCWNIQVNYIHKD NKSNGIVWCKHSMGFANMVNEGL QEQLYSLWESLASSANIDINIAK	None	None	None	None	None	None	None		

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